

CRF Errors Corrected by the STIC System Branch

0300 01P#6

Serial Number: 09/928,048A

CRF Processing Date: 5/16/02
 Edited by: _____
 Verified by: DC (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: **ENTERED**

- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____

- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____

- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____

- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file;
☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____

- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____

- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

***Examiner:** The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.



OIPE

RAW SEQUENCE LISTING

DATE: 05/16/2002

PATENT APPLICATION: US/09/928,048A

TIME: 13:00:53

Input Set : N:\jumbos\09928048A.DC.txt

Output Set: N:\CRF3\05162002\I928048A.raw

```

3 <110> APPLICANT: Scantibodies Laboratory, Inc.
4   Cantor, Thomas L.
6 <120> TITLE OF INVENTION: METHODS AND DEVICES FOR DIRECT
7   DETERMINATION OF CYCLASE INHIBITING PARATHYROID HORMONE
10 <130> FILE REFERENCE: 53221-20015.00
12 <140> CURRENT APPLICATION NUMBER: US 09/928,048A
13 <141> CURRENT FILING DATE: 2000-08-10
15 <160> NUMBER OF SEQ ID NOS: 8
17 <170> SOFTWARE: FastSEQ for Windows Version 4.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 9
21 <212> TYPE: PRT
22 <213> ORGANISM: Homo sapiens
24 <400> SEQUENCE: 1
25 Lys Lys Leu Gln Asp Val His Asn Phe
26 1          5
28 <210> SEQ ID NO: 2
29 <211> LENGTH: 5
30 <212> TYPE: PRT
31 <213> ORGANISM: Homo sapiens
33 <400> SEQUENCE: 2
34 Leu Gln Asp Val His
35 1          5
37 <210> SEQ ID NO: 3
38 <211> LENGTH: 84
39 <212> TYPE: PRT
40 <213> ORGANISM: Homo sapiens
42 <400> SEQUENCE: 3
43 Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn
44 1          5          10          15
45 Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His
46          20          25          30
47 Asn Phe Val Ala Leu Gly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser
48          35          40          45
49 Gln Arg Pro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu
50          50          55          60
51 Lys Ser Leu Gly Glu Ala Asn Lys Ala Asp Val Asn Val Leu Thr Lys
52 65          70          75          80
53 Ala Lys Ser Gln
56 <210> SEQ ID NO: 4
57 <211> LENGTH: 83
58 <212> TYPE: PRT
59 <213> ORGANISM: Homo sapiens

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61 <400> SEQUENCE: 4

62 Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn Ser

63 1 5 10 15

64 Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His Asn

65 20 25 30

66 Phe Val Ala Leu Gly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln

67 35 40 45

68 Arg Pro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu Lys

69 50 55 60

70 Ser Leu Gly Glu Ala Asn Lys Ala Asp Val Asn Val Leu Thr Lys Ala

71 65 70 75 80

72 Lys Ser Gln

75 <210> SEQ ID NO: 5

76 <211> LENGTH: 51

77 <212> TYPE: PRT

78 <213> ORGANISM: Homo sapiens

80 <400> SEQUENCE: 5

81 Phe Val Ala Leu Gly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln

82 1 5 10 15

83 Arg Pro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu Lys

84 20 25 30

85 Ser Leu Gly Glu Ala Asn Lys Ala Asp Val Asn Val Leu Thr Lys Ala

86 35 40 45

87 Lys Ser Gln

88 50

90 <210> SEQ ID NO: 6

91 <211> LENGTH: 34

92 <212> TYPE: PRT

93 <213> ORGANISM: Homo sapiens

95 <400> SEQUENCE: 6

96 Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn

97 1 5 10 15

98 Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His

99 20 25 30

100 Asn Phe

103 <210> SEQ ID NO: 7

104 <211> LENGTH: 50

105 <212> TYPE: PRT

106 <213> ORGANISM: Homo sapiens

108 <400> SEQUENCE: 7

109 Val Ala Leu Gly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln Arg

110 1 5 10 15

111 Pro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu Lys Ser

112 20 25 30

113 Leu Gly Glu Ala Asn Lys Ala Asp Val Asn Val Leu Thr Lys Ala Lys

114 35 40 45

115 Ser Gln

116 50

118 <210> SEQ ID NO: 8

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Input Set : N:\jumbos\09928048A.DC.txt

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119 <211> LENGTH: 20

120 <212> TYPE: PRT

121 <213> ORGANISM: Homo sapiens

123 <400> SEQUENCE: 8

124 Lys Ser Leu Gly Glu Ala Asn Lys Ala Asp Val Asn Val Leu Thr Lys

125 1 5 10 15

126 Ala Lys Ser Gln

127 20

VERIFICATION SUMMARY

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